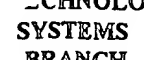


B. TECHNOLOGY SYSTEMS BRANCH



RECEIVED

MAY 16 2001

TECH CENTER 1600/2900,

Checker Version 3.0 can be down loaded from the USPTO website at the following address:
<http://www.uspto.gov/web/offices/pac/checker>

Raw Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/394,019

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

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MAY 16 2001

TECH CENTER 1600/2900

- 1 Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 2 Wrapped Aminos The amino acid number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 3 Incorrect Line Length The rules require that a line not exceed 72 characters in length. This includes spaces.
- 4 Misaligned Amino Acid The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs
Numbering between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
- 5 Non-ASCII This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
- 6 Variable Length Sequence(s) contain n's or Xaa's which represented more than one residue.
As per the rules, each n or Xaa can only represent a single residue.
Please present the maximum number of each residue having variable length and
indicate in the (ix) feature section that some may be missing.
- 7 PatentIn ver. 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid
sequence(s) . Normally, PatentIn would automatically generate this section from the
previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section
to the subsequent amino acid sequence. This applies primarily to the mandatory <220>-<223>
sections for Artificial or Unknown sequences.
- 8 Skipped Sequences Sequence(s) missing. If intentional, please use the following format for each skipped sequence:
(OLD RULES) (2) INFORMATION FOR SEQ ID NO:X:
 (i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:
 This sequence is intentionally skipped

Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
- 9 Skipped Sequences Sequence(s) missing. If intentional, please use the following format for each skipped sequence.
(NEW RULES) <210> sequence id number
 <400> sequence id number
 000
- 10 Use of n's or Xaa's Use of n's and/or Xaa's have been detected in the Sequence Listing.
(NEW RULES) Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 11 Use of "Artificial" Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules.
(NEW RULES) Valid response is Artificial Sequence.
- 12 Use of <220>Feature Sequence(s) are missing the <220>Feature and associated headings.
(NEW RULES) Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial Sequence" or "Unknown"
 Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
- 13 PatentIn ver. 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted
file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing).
Instead, please use "File Manager" or any other means to copy file to floppy disk.

1653

pp 1-5

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/394,019

DATE: 05/03/2001
TIME: 17:10:50

Input Set : A:\PTO.txt
Output Set: N:\CRF3\05032001\I394019.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: Oncoimmunin, Inc.
4 Komoriya, Akira
5 Packard, Beverly
7 <120> TITLE OF INVENTION: COMPOSITIONS FOR THE DETECTION OF ENZYME ACTIVITY IN BIOLOGICAL SAMPLES
8 AND ~~METHHODS~~ OF USE THEREOF **METHODS**
10 <130> FILE REFERENCE: M-903820US
12 <140> CURRENT APPLICATION NUMBER: US 09/394,019
13 <141> CURRENT FILING DATE: 1999-09-10
15 <150> PRIOR APPLICATION NUMBER: PCT/US98/00300
16 <151> PRIOR FILING DATE: 1998-02-20
18 <150> PRIOR APPLICATION NUMBER: US 08/802,981
19 <151> PRIOR FILING DATE: 1997-02-20
21 <160> NUMBER OF SEQ ID NOS: 257
23 <170> SOFTWARE: PatentIn version 3.0
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 12
27 <212> TYPE: PRT
C--> 28 <213> ORGANISM: Artificial/Unknown
30 <220> FEATURE:
31 <221> NAME/KEY: MOD_RES
32 <222> LOCATION: (3)..(3)
33 <223> OTHER INFORMATION: X is Aib or Pro
36 <400> SEQUENCE: 1
OK 38 Lys Asp Xaa Thr Gly Arg Thr Gly Pro Lys Gly Tyr
39 1 5 10
41 <210> SEQ ID NO: 2
42 <211> LENGTH: 14
43 <212> TYPE: PRT
C--> 44 <213> ORGANISM: Artificial/Unknown
46 <220> FEATURE:
W--> 47 <221> NAME/KEY: Artificial
48 <222> LOCATION: (1)..(14)
49 <223> OTHER INFORMATION: Artificial protease substrate
52 <400> SEQUENCE: 2
54 Lys Asp Pro Pro Thr Gly Arg Thr Gly Pro Pro Lys Gly Tyr
55 1 5 10
57 <210> SEQ ID NO: 3
58 <211> LENGTH: 14
59 <212> TYPE: PRT
C--> 60 <213> ORGANISM: Artificial/Unknown
62 <220> FEATURE:
63 <221> NAME/KEY: MOD_RES
64 <222> LOCATION: (3)..(3)
65 <223> OTHER INFORMATION: X is Aib or Pro
68 <400> SEQUENCE: 3
OK 70 Lys Asp Xaa Asp Glu Val Asp Gly Ile Asp Pro Lys Gly Tyr
71 1 5 10

invalid- Per 1.823 of Sequence Rules,
the only valid <213> responses are:

Unknown, Artificial Sequence,
or Scientific name
(Genus/species)

(one of the three)
do not combine

See circled portion
of item 12 on Euro
Summary Sheet

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/394,019

DATE: 05/03/2001
TIME: 17:10:50

Input Set : A:\PTO.txt
Output Set: N:\CRF3\05032001\I394019.raw

```

73 <210> SEQ ID NO: 4
74 <211> LENGTH: 14
75 <212> TYPE: PRT
C--> 76 <213> ORGANISM: Artificial/Unknown
78 <220> FEATURE:
79 <221> NAME/KEY: MOD_RES
80 <222> LOCATION: (3)..(3)
81 <223> OTHER INFORMATION: X is Aib or Pro
84 <400> SEQUENCE: 4
OK-> 86 Lys Asp Xaa Asp Glu Val Asn Gly Ile Asp Pro Lys Gly Tyr
87 1 5 10
89 <210> SEQ ID NO: 5
90 <211> LENGTH: 13
91 <212> TYPE: PRT
C--> 92 <213> ORGANISM: Artificial/Unknown
94 <220> FEATURE:
95 <221> NAME/KEY: MOD_RES
96 <222> LOCATION: (3)..(3)
97 <223> OTHER INFORMATION: X is Aib or Pro
100 <400> SEQUENCE: 5
OK-> 102 Lys Asp Xaa Glu Val Asp Gly Ile Asp Pro Lys Gly Tyr
103 1 5 10
105 <210> SEQ ID NO: 6
106 <211> LENGTH: 13
107 <212> TYPE: PRT
C--> 108 <213> ORGANISM: Artificial/Unknown
110 <220> FEATURE:
111 <221> NAME/KEY: MOD_RES
112 <222> LOCATION: (4)..(4)
113 <223> OTHER INFORMATION: X is Aib or Pro
116 <400> SEQUENCE: 6
OK-> 118 Lys Asp Tyr Xaa Ala Asp Gly Ile Asp Pro Lys Gly Tyr
119 1 5 10
121 <210> SEQ ID NO: 7
122 <211> LENGTH: 14
123 <212> TYPE: PRT
C--> 124 <213> ORGANISM: Artificial/Unknown
126 <220> FEATURE:
127 <221> NAME/KEY: MOD_RES
128 <222> LOCATION: (1)..(1)
129 <223> OTHER INFORMATION: BLOCKED WITH FMOC
132 <220> FEATURE:
133 <221> NAME/KEY: MOD_RES
134 <222> LOCATION: (3)..(3)
135 <223> OTHER INFORMATION: X is Aib or Pro
OK-> 138 <400> SEQUENCE: 7
140 Lys Asp Xaa Asp Glu Val Asp Gly Ile Asp Pro Lys Gly Tyr
141 1 5 10
143 <210> SEQ ID NO: 8

```

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/394,019

DATE: 05/03/2001
TIME: 17:10:50

Input Set : A:\PTO.txt
Output Set: N:\CRF3\05032001\I394019.raw

144 <211> LENGTH: 14
145 <212> TYPE: PRT
C--> 146 <213> ORGANISM: Artificial/Unknown
148 <220> FEATURE:
149 <221> NAME/KEY: MOD_RES
150 <222> LOCATION: (1)..(1)
151 <223> OTHER INFORMATION: BLOCKED with Fmoc
154 <220> FEATURE:
155 <221> NAME/KEY: MOD_RES
156 <222> LOCATION: (3)..(3)
157 <223> OTHER INFORMATION: X is Aib or Pro
160 <400> SEQUENCE: 8
OK> 162 Lys Asp Xaa Asp Glu Val Asn Gly Ile Asp Pro Lys Gly Tyr
163 1 5 10
165 <210> SEQ ID NO: 9
166 <211> LENGTH: 13
167 <212> TYPE: PRT
C--> 168 <213> ORGANISM: Artificial/Unknown
170 <220> FEATURE:
171 <221> NAME/KEY: MOD_RES
172 <222> LOCATION: (1)..(1)
173 <223> OTHER INFORMATION: BLOCKED with Fmoc
176 <220> FEATURE:
177 <221> NAME/KEY: MOD_RES
178 <222> LOCATION: (3)..(3)
179 <223> OTHER INFORMATION: X is Aib or Pro
182 <400> SEQUENCE: 9
OK> 184 Lys Asp Xaa Glu Val Asp Gly Ile Asp Pro Lys Gly Tyr
185 1 5 10
187 <210> SEQ ID NO: 10
188 <211> LENGTH: 12
189 <212> TYPE: PRT
C--> 190 <213> ORGANISM: Artificial/Unknown
192 <220> FEATURE:
193 <221> NAME/KEY: MOD_RES
194 <222> LOCATION: (1)..(1)
195 <223> OTHER INFORMATION: BLOCKED with Fmoc
198 <400> SEQUENCE: 10
200 Lys Asp Ala Ile Pro Met Ser Ile Pro Lys Gly Tyr
201 1 5 10
203 <210> SEQ ID NO: 11
204 <211> LENGTH: 12
205 <212> TYPE: PRT
C--> 206 <213> ORGANISM: Artificial/Unknown
208 <220> FEATURE:
209 <221> NAME/KEY: MOD_RES
210 <222> LOCATION: (1)..(1)
211 <223> OTHER INFORMATION: BLOCKED with Fmoc
214 <220> FEATURE:

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/394,019

DATE: 05/03/2001
TIME: 17:10:50

Input Set : A:\PTO.txt
Output Set: N:\CRF3\05032001\I394019.raw

215 <221> NAME/KEY: MOD_RES
216 <222> LOCATION: (6)..(6)
217 <223> OTHER INFORMATION: X is norleucine
220 <400> SEQUENCE: 11
OK 222 Lys Asp Ala Ile Pro Xaa Ser Ile Pro Lys Gly Tyr
223 1 5 10
225 <210> SEQ ID NO: 12
226 <211> LENGTH: 16
227 <212> TYPE: PRT
C--> 228 <213> ORGANISM: Artificial/Unknown
230 <220> FEATURE:
231 <221> NAME/KEY: MOD_RES
232 <222> LOCATION: (1)..(1)
233 <223> OTHER INFORMATION: BLOCKED with Fmoc
236 <220> FEATURE:
237 <221> NAME/KEY: MOD_RES
238 <222> LOCATION: (3)..(3)
239 <223> OTHER INFORMATION: X is Aib or Pro
242 <400> SEQUENCE: 12
OK 244 Lys Asp Xaa Gly Asp Glu Val Asp Gly Ile Asp Gly Pro Lys Gly Tyr
245 1 5 10 15
247 <210> SEQ ID NO: 13
248 <211> LENGTH: 18
249 <212> TYPE: PRT
C--> 250 <213> ORGANISM: Artificial/Unknown
252 <220> FEATURE:
253 <221> NAME/KEY: MOD_RES
254 <222> LOCATION: (1)..(1)
255 <223> OTHER INFORMATION: BLOCKED with Fmoc
258 <220> FEATURE:
259 <221> NAME/KEY: MOD_RES
260 <222> LOCATION: (14)..(14)
261 <223> OTHER INFORMATION: X is epsilon aminocaproic acid
264 <220> FEATURE:
265 <221> NAME/KEY: MOD_RES
266 <222> LOCATION: (4)..(4)
267 <223> OTHER INFORMATION: X is epsilon aminocaproic acid
270 <220> FEATURE:
271 <221> NAME/KEY: MOD_RES
272 <222> LOCATION: (3)..(3)
273 <223> OTHER INFORMATION: X is Aib or Pro
276 <400> SEQUENCE: 13
OK 278 Lys Asp Xaa Xaa Gly Asp Glu Val Asp Gly Ile Asp Gly Xaa Pro Lys
279 1 5 10 15
281 Gly Tyr
284 <210> SEQ ID NO: 14
285 <211> LENGTH: 18
286 <212> TYPE: PRT
C--> 287 <213> ORGANISM: Artificial/Unknown

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/394,019

DATE: 05/03/2001
TIME: 17:10:50

Input Set : A:\PTO.txt
Output Set: N:\CRF3\05032001\I394019.raw

289 <220> FEATURE:
290 <221> NAME/KEY: MOD_RES
291 <222> LOCATION: (1)..(1)
292 <223> OTHER INFORMATION: MODIFIED WITH benzyloxycarbonyl
295 <220> FEATURE:
296 <221> NAME/KEY: MOD_RES
297 <222> LOCATION: (14)..(14)
298 <223> OTHER INFORMATION: X is epsilon aminoacproic acid
301 <220> FEATURE:
302 <221> NAME/KEY: MOD_RES
303 <222> LOCATION: (3)..(3)
304 <223> OTHER INFORMATION: X is Aib or Pro
307 <220> FEATURE:
308 <221> NAME/KEY: MOD_RES
309 <222> LOCATION: (4)..(4)
310 <223> OTHER INFORMATION: X is epsilon aminoacproic acid
313 <400> SEQUENCE: 14
315 Lys Asp Xaa Xaa Gly Asp Glu Val Asp Gly Ile Asp Gly Xaa Pro Lys
316 1, . . . 5 10 15
318 Gly Tyr
321 <210> SEQ ID NO: 15
322 <211> LENGTH: 13
323 <212> TYPE: PRT
C--> 324 <213> ORGANISM: Artificial/Unknown
326 <220> FEATURE:
327 <221> NAME/KEY: MOD_RES
328 <222> LOCATION: (1)..(1)
329 <223> OTHER INFORMATION: BLOCKED with Fmoc
332 <400> SEQUENCE: 15
334 Lys Asp Tyr Asx Ala Asp Gly Ile Asp Pro Lys Gly Tyr
335 1 . . . 5 10
337 <210> SEQ ID NO: 16
338 <211> LENGTH: 18
339 <212> TYPE: PRT
340 <213> ORGANISM: artificial sequence
342 <220> FEATURE:
343 <221> NAME/KEY: MOD_RES
344 <222> LOCATION: (1)..(1)
345 <223> OTHER INFORMATION: Blocked with Fa
348 <220> FEATURE:
349 <221> NAME/KEY: MOD_RES
350 <222> LOCATION: (4)..(4)
351 <223> OTHER INFORMATION: X is epsilon amino caproic acid
354 <400> SEQUENCE: 16
W--> 356 Lys Asp Pro Xaa Gly Asp Glu Val Asp Gly Ile Asn Gly Xaa Pro Lys
357 1 . . . 5 10 15
359 Gly Tyr
362 <210> SEQ ID NO: 17
363 <211> LENGTH: 16

global error (see below)

*what about
this Xaa?*

*the only
2207-2237
explanation
is for Xaa
at
location
4*

The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

Please Note:

Use f n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields f each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/394,019

DATE: 05/03/2001
TIME: 17:10:51

Input Set : A:\PTO.txt
Output Set: N:\CRF3\05032001\I394019.raw

L:28 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:38 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:44 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
L:47 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2
L:60 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:70 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:76 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:86 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:92 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:102 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:108 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:118 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:124 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:140 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:146 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
L:162 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:168 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:184 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:190 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:206 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
L:222 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:228 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:244 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:250 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13
L:278 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:287 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:315 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:324 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
L:356 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:387 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:415 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:440 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:459 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:490 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:512 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:537 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:562 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:587 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:612 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:643 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:680 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:711 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:736 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:761 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:786 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:811 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:836 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
L:861 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/394,019

DATE: 05/03/2001
TIME: 17:10:51

Input Set : A:\PTO.txt
Output Set: N:\CRF3\05032001\I394019.raw

L:898 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36
L:935 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37
L:966 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38
L:1003 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39
L:1040 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40
L:1077 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41
L:1096 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
L:1112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43
L:1128 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44
L:1144 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45
L:1160 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46
L:1176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47
L:1204 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48
L:1235 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49
L:1266 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50
L:1297 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:1334 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:1365 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53
L:3524 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:128
L:3530 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:128
L:3536 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:128
L:4385 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:161
L:4388 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:161
L:4407 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:162
L:4420 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:163
L:4423 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:163
L:4436 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:164
L:4452 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:165
L:4468 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:166
L:4496 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:167
L:4524 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:168
L:4564 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:169
L:4598 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:170
L:4632 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:171
L:4666 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:172
L:4694 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:173
L:4728 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:174
L:4756 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:175
L:4784 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:176
L:4812 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:177
L:4846 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:178
L:4895 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:179
L:4944 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:180
L:4978 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:181
L:5006 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:182
L:5046 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:183
L:5095 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:184
L:5144 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:185
L:5172 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:186

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/394,019

DATE: 05/03/2001
TIME: 17:10:51

Input Set : A:\PTO.txt

Output Set: N:\CRF3\05032001\I394019.raw

L:5188 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:187
L:5210 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:188
L:5238 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:189
L:5260 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:190
L:5294 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:191
L:5328 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:192
L:5377 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:193
L:5380 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:193
L:5393 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:194
L:5409 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:195
L:5412 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:195
L:5425 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:196
L:5428 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:196
L:5444 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:197
L:5460 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:198
L:5476 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:199
L:5492 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:200
L:5536 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:202
L:5552 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:203
L:5606 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:206
L:5622 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:207
L:5638 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:208
L:5654 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:209
L:5670 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:210
L:5686 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:211
L:5702 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:212
L:5718 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:213
L:5734 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:214
L:5750 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:215
L:5794 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:217
L:5810 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:218
L:5826 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:219